hw1

January 24, 2024

```
[25]: import sys
      if sys.version_info[0] < 3:</pre>
         raise Exception("Python 3 not detected.")
      import numpy as np
      import matplotlib.pyplot as plt
      from sklearn import svm
      from scipy import io
      if __name__ == "__main__":
          for data_name in ["mnist", "spam", "toy"]:
             data = np.load(f"data/{data_name}-data.npz")
             print("\nloaded %s data!" % data_name)
             fields = "test_data", "training_data", "training_labels"
             for field in fields:
                print(field, data[field].shape)
     loaded mnist data!
     test_data (10000, 1, 28, 28)
     training_data (60000, 1, 28, 28)
     training_labels (60000,)
     loaded spam data!
     test_data (1000, 39)
     training_data (4171, 39)
     training_labels (4171,)
     loaded toy data!
     test_data (0,)
     training_data (1000, 2)
     training_labels (1000,)
     0.1 3. Data Partition
[26]: np.random.seed(15)
      """Shuffles and partitions data"""
```

```
def partition(data, labels, validation_size):
    total_size = len(data)
    # in the case where a percentage is given
    if validation_size < 1:</pre>
        validation_size = int(validation_size * total_size)
    shuffled_ind = np.random.permutation(total_size)
    # uses fancy indexing, first reshuffling data, then getting the validation_
 ⇔set
    val_data = data[shuffled_ind][:validation_size]
    val_label = labels[shuffled_ind][:validation_size]
    train_data = data[shuffled_ind][validation_size:]
    train_label = labels[shuffled_ind][validation_size:]
    return train_data, train_label, val_data, val_label
def eval_metric(y_pred, y_hat):
    assert len(y_pred) == len(y_hat)
    total_pred = len(y_pred)
    correct_pred = np.sum(y_pred == y_hat) # vectorized, easier than for loop
    return correct_pred / total_pred
```

```
[27]: mnist = np.load(f"data/mnist-data.npz")
      spam = np.load(f"data/spam-data.npz")
      mnist_train_data_flat = mnist["training_data"].reshape(
          mnist["training_data"].shape[0], -1
      )
      minst_train_d, minst_train_l, minst_val_d, minst_val_l = partition(
          mnist_train_data_flat, mnist["training_labels"], 10000
      )
      spam_train_d, spam_train_l, spam_val_d, spam_val_l = partition(
          spam["training_data"], spam["training_labels"], 0.2
      )
      print(
          len(minst_train_l),
          len(minst_val_1),
          len(minst_val_d),
          len(spam_train_1),
          len(spam_train_d),
          len(spam_val_1),
      )
```

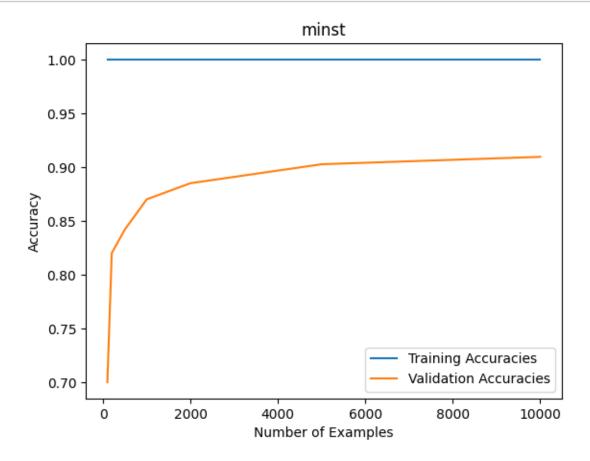
50000 10000 10000 3337 3337 834

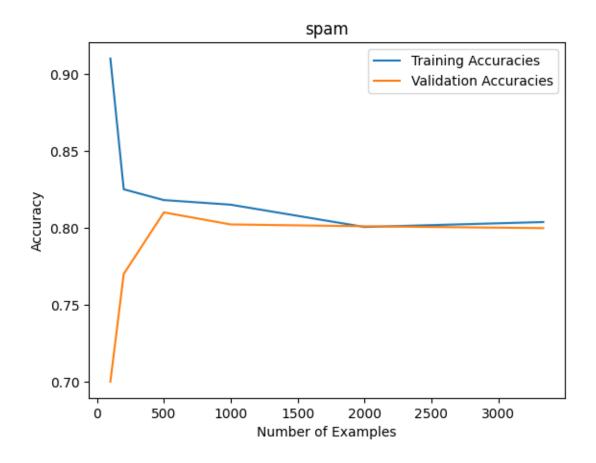
0.2 4. Support Vector Machines

```
[16]: def svm_model(data, training_sizes, train_data, train_label, val_data,_
       ⇔val_label):
          # store accuracies across diff training sizes
          train_accuracies = []
          val accuracies = []
          for size in training sizes:
              # Train model
              model = svm.SVC(kernel="linear")
              model.fit(train_data[:size], train_label[:size])
              # Predict training accuracy
              train_pred = model.predict(train_data[:size])
              train_accuracy = eval_metric(train_pred, train_label[:size])
              train_accuracies.append(train_accuracy)
              # Predict validation accuracy
              val_pred = model.predict(val_data[:size])
              val_accuracy = eval_metric(val_pred, val_label[:size])
              val_accuracies.append(val_accuracy)
          plot_ex_accuracies(data, training_sizes, train_accuracies, val_accuracies)
      def plot_ex_accuracies(data, training_sizes, train_accuracies, val_accuracies):
         plt.figure()
          plt.plot(training_sizes, train_accuracies, label="Training Accuracies")
          plt.plot(training_sizes, val_accuracies, label="Validation Accuracies")
          plt.title(data)
          plt.xlabel("Number of Examples")
          plt.ylabel("Accuracy")
          plt.legend()
[46]: m_train_sizes = [100, 200, 500, 1000, 2000, 5000, 10000]
      s_train sizes = [100, 200, 500, 1000, 2000, spam_train d.shape[0]]
      # print("loaded mnist data!")
      # fields = "test_data", "training_data", "training_labels"
      # for field in fields:
           print(field, mnist[field].shape)
      # print(mnist_train_data_flat.shape)
      svm_model(
          "minst", m_train_sizes, minst_train_d, minst_train_l, minst_val_d,

→minst_val_l
```

```
)
svm_model("spam", s_train_sizes, spam_train_d, spam_train_l, spam_val_d,u
spam_val_l)
```





0.3 5. Hyperparameter Tuning

I tried these C values [0.00000001, 0.0000001, 0.000001, 0.00001, 0.0001, 0.0001, 0.001, 0.01, 0.05, 0.1, 1], the corresponding accuracy was 1e-08:0.8883 1e-07:0.9222 1e-06:0.9298 1e-05:0.918 0.0001:0.9095 0.001:0.9095 0.01:0.9095 0.05:0.9095 0.1:0.9095 1:0.9095

1e-08: 0.8883
1e-07: 0.9222
1e-06: 0.9298
1e-05: 0.918
0.0001: 0.9095
0.001: 0.9095
0.05: 0.9095
0.1: 0.9095
1: 0.9095

0.4 6. K-fold cross validation

I tried the C values listbed below and these were the accuracies, the best C value is 10 in my case C=1e-05: Cross-validation accuracy = 0.7123028762618647 C=0.0001: Cross-validation accuracy = 0.7168577952009649 C=0.001: Cross-validation accuracy = 0.7487428021654533 C=0.01: Cross-validation accuracy = 0.7753540401211965 C=0.05: Cross-validation accuracy = 0.791898217952584 C=0.1: Cross-validation accuracy = 0.7926170680222289 C=1: Cross-validation accuracy = 0.8005293011100102 C=10: Cross-validation accuracy = 0.8036456583236407 C=50: Cross-validation accuracy = 0.8029265210586022 C=100: Cross-validation accuracy = 0.8034061373655568

```
[18]: def k fold_cross_validation(train_data, train_label, c_values, k=5):
          total_size = len(train_data)
          shuffled_ind = np.random.permutation(total_size)
          fold_size = total_size // k
          C_accuracies = []
          for c in c_values:
              fold accuracies = []
              for fold in range(k):
                  # indices for val set
                  start, end = (
                      fold * fold_size,
                      (fold + 1) * fold_size if fold < k - 1 else total_size,
                  )
                  # val, training indices
                  val_indices = shuffled_ind[start:end]
                  train_indices = np.concatenate((shuffled_ind[:start],__
       ⇒shuffled ind[end:]))
                  # slice data
                  train_fold_data, train_fold_label = (
                      train data[train indices],
                      train_label[train_indices],
                  val_fold_data, val_fold_label = (
                      train_data[val_indices],
                      train_label[val_indices],
                  )
                  k_fold_model = svm.SVC(kernel="linear", C=c)
                  k_fold_model.fit(train_fold_data, train_fold_label)
                  val_predict = k_fold_model.predict(val_fold_data)
                  val_accuracy = eval_metric(val_predict, val_fold_label)
                  fold_accuracies.append(val_accuracy)
              avg_accuracy = np.mean(fold_accuracies)
```

```
C_accuracies.append((c, avg_accuracy))
             print(f"C={c}: Cross-validation accuracy = {avg_accuracy}")
         C_accuracies.sort(
             key=lambda x: x[1], reverse=True
         ) # sort by descending avg_accuracy
         return C_accuracies
[7]: spam = np.load(f"data/spam-data.npz")
     k fold cross validation(
         spam["training_data"],
         spam["training_labels"],
         [0.00001, 0.0001, 0.001, 0.01, 0.05, 0.1, 1, 10, 50, 100],
     )
    C=1e-05: Cross-validation accuracy = 0.7123028762618647
    C=0.0001: Cross-validation accuracy = 0.7168577952009649
    C=0.001: Cross-validation accuracy = 0.7487428021654533
    C=0.01: Cross-validation accuracy = 0.7753540401211965
    C=0.05: Cross-validation accuracy = 0.791898217952584
    C=0.1: Cross-validation accuracy = 0.7926170680222289
    C=1: Cross-validation accuracy = 0.8005293011100102
    C=10: Cross-validation accuracy = 0.8036456583236407
    C=50: Cross-validation accuracy = 0.8029265210586022
    C=100: Cross-validation accuracy = 0.8034061373655568
    C=1000: Cross-validation accuracy = 0.8036459455190339
[7]: [(1000, 0.8036459455190339),
      (10, 0.8036456583236407),
      (100, 0.8034061373655568),
      (50, 0.8029265210586022),
      (1, 0.8005293011100102),
      (0.1, 0.7926170680222289),
      (0.05, 0.791898217952584),
      (0.01, 0.7753540401211965),
      (0.001, 0.7487428021654533),
      (0.0001, 0.7168577952009649),
      (1e-05, 0.7123028762618647)]
```

0.5 Question 7: Kaggle Submissions

My submission on Kaggle had a score of Mnist: 0.978, and Spam: 0.84. For both datasets, I wrote a function that finds the best model and hyperparameters for me, experiementing on different kernels: poly, rbf; different values of C, different values of gamma and so on. By sorting though this I was able to find models for mnist that worked really well using "poly", C = 0.000001, gamma = 0.01. However, there doesn't seem to be too much of an improvement in Spam, especially since k-fold modeling wasn't available (can only submit 1 model). So for Spam, I read through some of the emails and experimented with adding features such as "credit", "\$", "offer", and used my general intuition on spam/ham emails to add even more features like "click", "urgent". By adding suich features and tuning hyperparameters and using rbf, I was able to achieve a 84% score for spam.

```
[32]: def find_best_model(dataset, train_data, train_label, val_data, val_label, c):
          val_accuracies = []
          gammas = [0.001, 0.01, 0.1]
          kernels = ["poly", "rbf"]
          for kernel in kernels:
              for gamma in gammas:
                  val_accuracy = train_model(
                       dataset,
                       train_data,
                       train label,
                       val data,
                       val_label,
                       kernel,
                       С,
                       gamma,
                  val_accuracies.append((val_accuracy, kernel, c, gamma))
                  print(f"Accuracy: {val_accuracy}, kernel: {kernel}, C={c}, gamma:__
       →{gamma}")
          return val_accuracies.sort(key=lambda x: x[1], reverse=True)
      def train model(
          dataset, train_data, train_label, val_data, val_label, kernel, c, gamma
      ):
          # if dataset == "spam":
                accuracy = k fold cross_validation(kernel, c, gamma, train_data,__
       \hookrightarrow train\_label)
                return accuracy
          # else:
              model = svm.SVC(kernel=kernel, C=c, gamma=gamma)
              model = model.fit(train_data, train_label)
              val_predict = model.predict(val_data)
              val_accuracy = eval_metric(val_predict, val_label)
              return val_accuracy
```

```
# def k fold cross_validation(kernel, c, qamma, train_data, train_label, k=5):
      total \ size = len(train \ data)
      shuffled_ind = np.random.permutation(total_size)
     fold_size = total_size // k
#
#
     fold_accuracies = []
#
      for fold in range(k):
#
          # indices for val set
#
          start, end = (
              fold * fold_size,
#
              (fold + 1) * fold_size if fold < k - 1 else total_size,
#
#
          # val, training indices
#
          val_indices = shuffled_ind[start:end]
          train_indices = np.concatenate((shuffled_ind[:start],_
 →shuffled_ind[end:]))
          # slice data
#
          train fold data, train fold label = (
              train_data[train_indices],
              train_label[train_indices],
#
#
          val_fold_data, val_fold_label = (
#
              train_data[val_indices],
              train_label[val_indices],
#
          k_fold_model = svm.SVC(kernel=kernel, C=c, qamma=qamma)
#
          k fold model.fit(train data, train label)
          val\_predict = k\_fold\_model.predict(val\_fold\_data)
#
          val_accuracy = eval_metric(val_predict, val_fold_label)
#
          fold_accuracies.append(val_accuracy)
#
      avg_accuracy = np.mean(fold_accuracies)
      return avg_accuracy
```

```
spam_train_d, spam_train_l, spam_val_d, spam_val_l = partition(
          spam["training_data"], spam["training_labels"], 0.2
[36]: find_best_model(
          "spam", spam_train_d, spam_train_l, spam_val_d, spam_val_l, 10
      ) # Accuracy: 0.8759328855751498, kernel: rbf, C=10, gamma: 0.1
     Accuracy: 0.6990407673860911, kernel: poly, C=10, gamma: 0.001
     Accuracy: 0.7278177458033573, kernel: poly, C=10, gamma: 0.01
     Accuracy: 0.7649880095923262, kernel: poly, C=10, gamma: 0.1
     Accuracy: 0.7685851318944844, kernel: rbf, C=10, gamma: 0.001
     Accuracy: 0.8321342925659473, kernel: rbf, C=10, gamma: 0.01
     Accuracy: 0.841726618705036, kernel: rbf, C=10, gamma: 0.1
[20]: find_best_model(
          "minst", mnist_train_d, mnist_train_l, mnist_val_d, mnist_val_l, 0.000001
       # Accuracy: 0.9789, kernel: poly, C=1e-06, gamma: 0.01
     Accuracy: 0.9775, kernel: poly, C=1e-06, gamma: 0.001
     Accuracy: 0.9775, kernel: poly, C=1e-06, gamma: 0.01
     Accuracy: 0.9775, kernel: poly, C=1e-06, gamma: 0.1
     0.5.1 Kaggle
[39]: import pandas as pd
      def kaggle_submit(dataset, train_data, train_labels, test_data, kernel, c,_
       ⇒gamma):
          model = svm.SVC(kernel=kernel, C=c, gamma=gamma)
          model.fit(train data, train labels)
          if dataset == "mnist":
              test_data = test_data.reshape(test_data.shape[0], -1)
          test_labels = model.predict(test_data)
          test_labels = test_labels.astype(int)
          df = pd.DataFrame({"Category": test_labels})
          df.index += 1
          df.to_csv(f"{dataset}_predictions.csv", index_label="Id")
[40]: kaggle_submit("spam", spam_train_d, spam_train_l, spam["test_data"], "rbf", 10,
       →0.1)
      # kaggle submit(
            "mnist", mnist_train_d, mnist_train_l, mnist["test_data"], "poly", 0.
       →000001, 0.01
```

)
[]: